# Making the Most of What We Know: Towards Effective Use of Genomics Data

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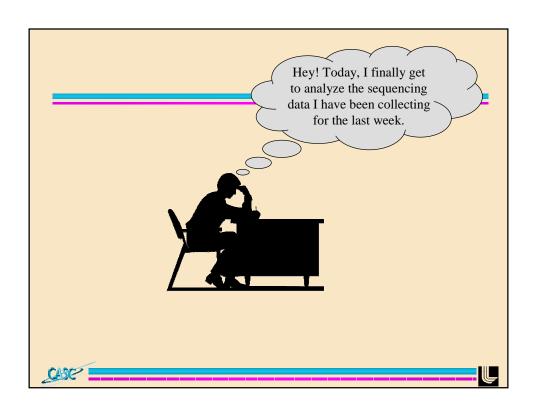
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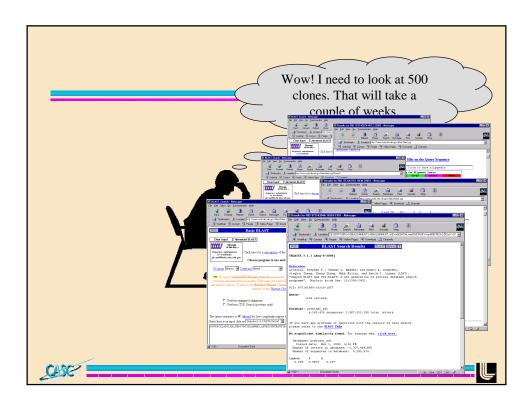


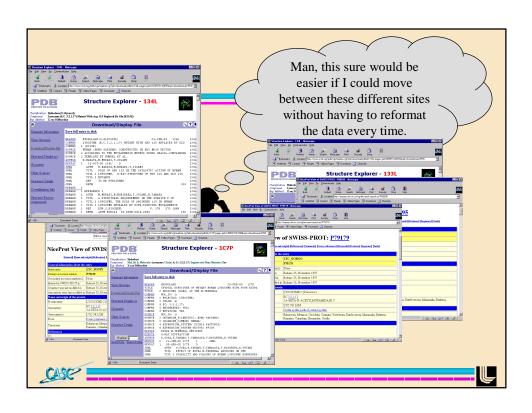
### **Outline**

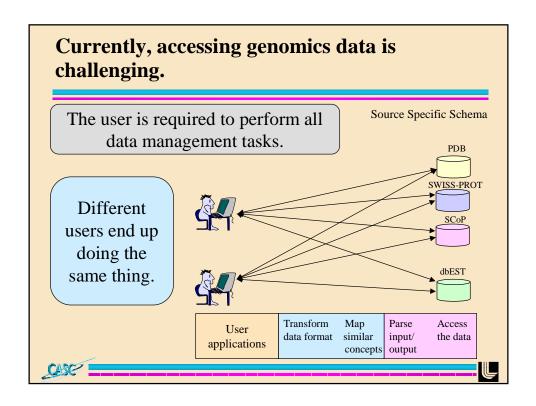
- A day in the life of a geneticist/biologist.
- The business approach. (present)
- The impact of the web. (future)
- The key to success.

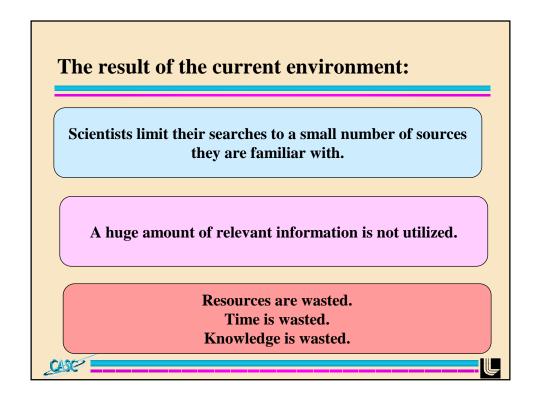


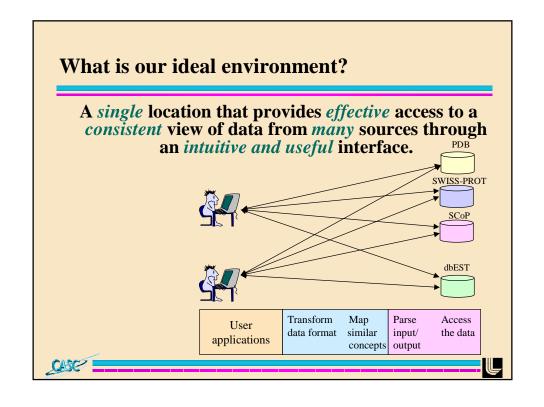


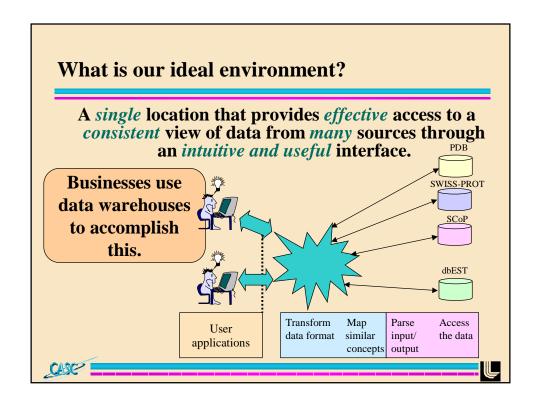


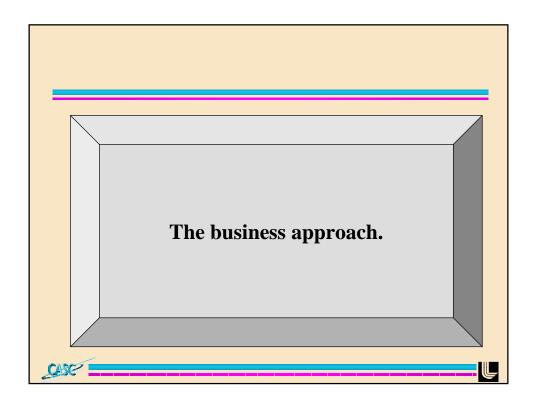


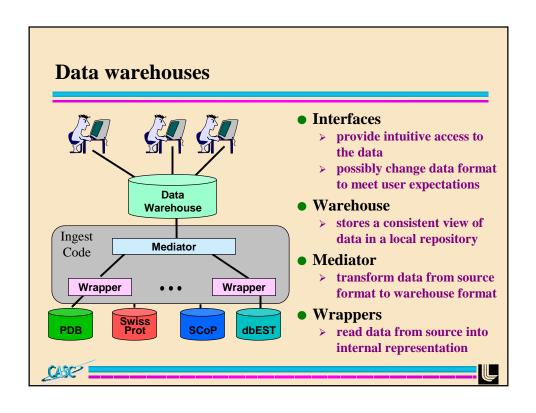


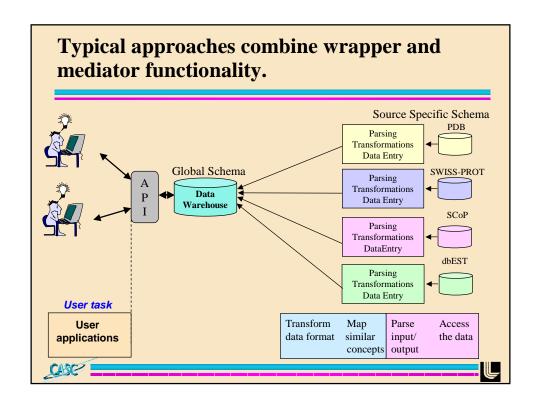


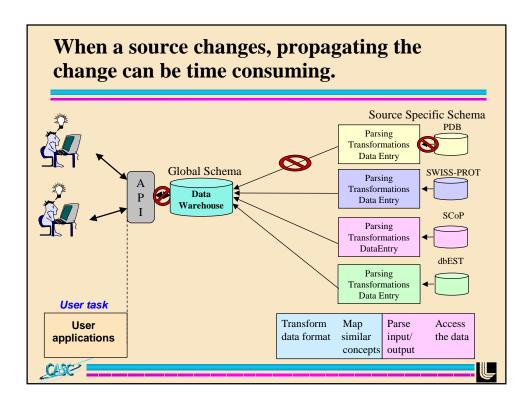


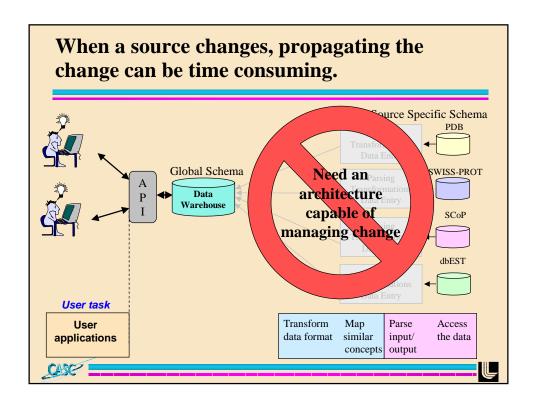


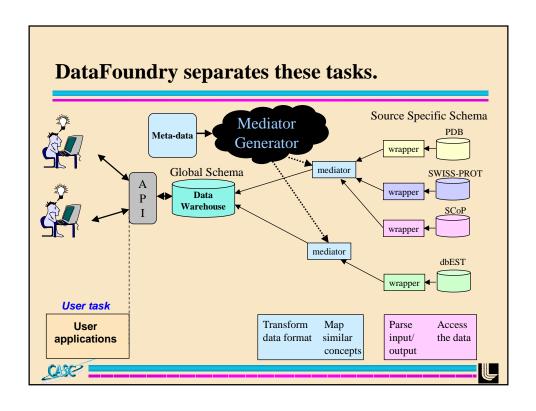


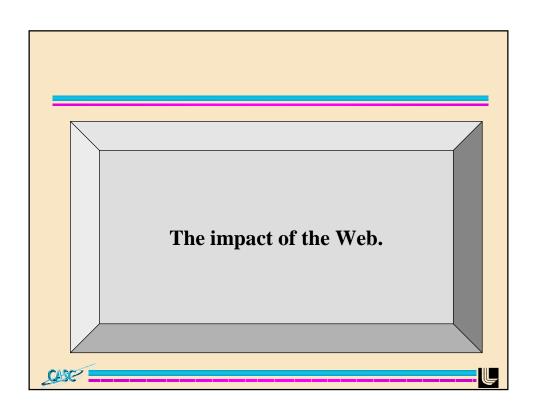


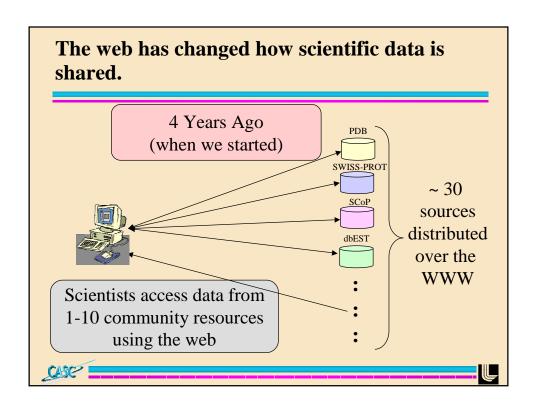


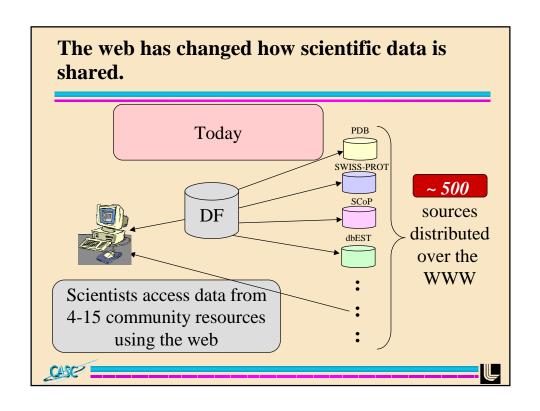










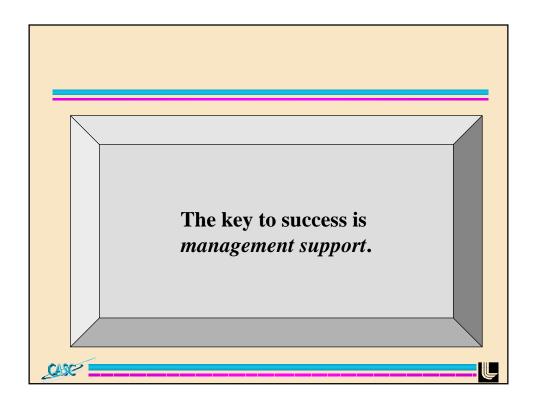


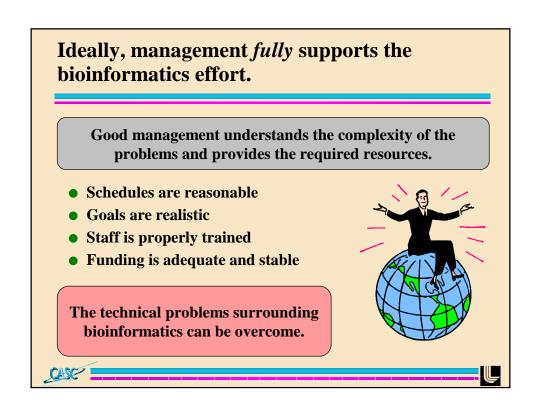
## So, what makes this so much more difficult?

- No central directory
  - > How do you find the sources in the first place?
- Different data formats and semantics
  - > Once you find a source, how to you make sense of it?
- Complex analysis
  - > Queries require more than simple data retrieval, they need to invoke complex programs.
- Lots of data sources
  - > Too much work to be done manually.
  - > Need to select appropriate subset for each user / query.
  - > How do you present the results in an useful format?



# A hybrid approach to information access holds the most promise. Source Specific Schema Global Schema SWISS-PROT Detailed analysis Dynamic Ouerv dbEST source Engine access Meta-data to large numbers of non-integrated **Preprocess**





## Unfortunately, that is usually not the case.

Poor management underestimates the complexity of the problems and provides inadequate resources.

- Schedules are unreasonable
- Goals are unrealistic
- Morale is low & staff is poorly trained,
- Funding is minimal and unstable

You are contributing to the problems facing bioinformatics instead of their solutions.





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# **Conclusions**

Bioinformatics has a long way to go before it can support scientific exploration.

#### The good news

The technical challenges surrounding bioinformatics can be overcome.

#### The bad news

The major challenge is political, not technical.



